1651

## **Cephalosporin C Biosynthesis; a Branched Pathway Sensitive to a Kinetic Isotope Effect**

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Incubation of [3-2H] penicillin N with preparations of deacetoxycephalosporin C/deacetylcephalosporin C synthetase activity from *Cephalosporium acremonium* CO *728* gave, along with the normal product deacetoxycephalosporin C, another β-lactam metabolite, namely 7β-[(R)-5-amino-5-carboxypentanoyl]-3β-hydroxy-3α-methyl[4-2H)-cepham- $4\alpha$ -carboxylic acid. This material arises as a result of a deuterium isotope effect on a branched pathway in the enzymic mechanism. The 3B-hydroxy group in this substance arises from molecular oxygen.

Although the conversion of penicillin N (1) into deacetoxyce-<br>
phalosporin C (DAOC) (2) and deacetylcephalosporin C pentanoyl]-3β-hydroxy-3α-methylcepham-4α-carboxylic acid phalosporin C **(DAOC) <sup>(2</sup>)** and deacetylcephalosporin C **pentanoyl**-3 $\beta$ -hydroxy-3 $\alpha$ -methylcepham-4 $\alpha$ -carboxylic acid **(DAC)** (3) by a cell-free extract from *Cephalosporium* **(4a)**† from a filtered broth of *C*. *a acremonium* was first rigorously proven in 1980,<sup>1</sup> no inter- mechanistic speculation. The involvement of an episulphomediates in this process have been subsequently revealed (Scheme 1). Both sequential steps consume dioxygen and  $+$  We have previously referred to the  $(R)$ -5-amino-5-carboxypentan-<br> $\alpha$ -ketoglutarate, and the latter is converted into succinate and oyl group as  $\delta$ -(p- $\alpha$ -amin a-ketoglutarate, and the latter is converted into succinate and

**(4a)**<sup> $\dagger$ </sup> from a filtered broth of *C. acremonium* led to early



**<sup>a</sup>**DAC **(3)** was treated with formic acid to give the lactone **(6)** prior to mass spectral analysis. Samples were then run under positive ion thermospray h.p.1.c. mass spectrometry conditions, using a reverse phase octadecylsilane column with 0.05M-ammonium acetate containing 1% acetonitrile adjusted to pH *5* with formic acid as eluant. **b** Samples evaporated on stage from water, pre-protonated with 5% methanolic oxalic acid, then run under positive argon fast atom bombardment using glycerol as matrix. *c* For synthetic sample of (4a). <sup>d</sup> Calculated for  $C_{14}H_{22}N_3O_7S$ 



nium ion *(5)* was suggested, which could directly collapse *via*  proton loss to **(2),** or alternatively be intercepted by water to give (4a), (Scheme 2).<sup>2</sup> More recently, chemical modelling of the ring-expansion step has led to the suggestion that an

equilibrating free radical (Scheme 3) could equally well explain this reaction,3 but clearly heterolytic trapping by water of such a radical to give **(4a)** would be unreasonable. We now report the results of using [3-2H]penicillin N **(lb)** on the course **of** the enzymic reaction.

Initially the product composition of an incubation of penicillin N **(la)** with partially purified **DAOC/DAC** synthe-



tase from *C. acremonium* CO 728‡ and standard co-factors§ was examined by 500 MHz n.m.r. and shown to contain three products  $[(2) : (3) : (4a) \ 40 : 20 : 1]$ . Repeating the incubation with [3-<sup>2</sup>H]penicillin N (1b)] gave the same three products *but in substantially different ratio,* **[(2)** : **(3)** : **(4b)** 40 : 25 : 351.

The structure of **(4b),** purified by h.p.1.c. [reverse phase octadecylsilane column; 25 mm-NH<sub>4</sub>HCO<sub>3</sub> as eluant], was determined from its spectral data (consistent with literature values)<sup>2</sup> and by chemical synthesis<sup>5</sup> of **(4a)**. For **(4b)**  $\delta_H$  (500 MHz; D<sub>2</sub>O, ref. sodium 3-trimethylsilyl [2,2,3,3-<sup>2</sup>H<sub>4</sub>] propanoate) 1.38(3H, s, Me), 1.65-1.80 and 1.85-1.95(4H,  $2 \times$ m,  $(CH<sub>2</sub>)<sub>2</sub>CH<sub>2</sub>CO$ , 2.42(2H, *ca.* t, *J* 7.5 Hz, CH<sub>2</sub>CO), 2.64 and 3.54(2H, ABq, *J* 14 Hz, 2-H), 3.6-3.7(1H, m, CH[CH<sub>2</sub>]<sub>3</sub>), and 5.29 and 5.44 (2  $\times$  1H, 2  $\times$  d, J 4 Hz, 6-H, 7-H); *m/z* (positive argon fast atom bombardment) 377 *(MH+);* no antibacterial activity towards *Staphyfococcus aureus* N.C.T.C. 6571 or *Escherichia cofi ESS* at a concentration of  $100 \mu g$  ml<sup>-1</sup> (sample size  $100 \mu l$ ). The hydroxycepham **(4b)** was shown *not* to be a substrate for cephem formation with DAOC/DAC synthetase in separate experiments.

Secondly, we examined the origin of the  $3\beta$ -hydroxy function of **(4).** Thus incubation of **(lb)** under a closed atmosphere of  ${}^{18}O_2$  gas (99%) gave both labelled 3 $\beta$ -hydroxycepham **(4b)** and DAC **(3),** which were purified by h.p.1.c. Lactonisation of **(3)** to **(6)** (formic acid) gave a sample suitable for mass spectral analysis. This technique revealed <sup>18</sup>O incorporation into both  $(3)^6$  and the  $3\beta$ -hydroxycepham **(4b)\*\*** (Table 1). **A** similar analysis of the co-produced DAOC  $(2)$  from the  ${}^{18}O_2$  experiment revealed *no* label incorporation.



These experiments require that the conversion of penicillin N **(la)** into **(2)** proceeds *via* a branched pathway through an intermediate which provides (2) as well as (4).<sup>††</sup> With unlabelled penicillin N (1a) the ratio  $(2) + (3) : (4)$  is 60 : 1, so that **(4)** is **a** minor product of the ring-expansion step, explaining its low concentration relative to cephalosporin C in normal fermentations.<sup>2</sup> However the operation of a deuterium isotope effect on the breakage of the C(3)-H bond (penicillin numbering) substantially shifts the above ratio to *ca.* 2 : 1. Additionally, the hydroxy-containing product **(4)** is formed by the specific incorporation of oxygen from *dioxygen.*  All these facts may be accommodated by a mechanism (Scheme 4), in which a bridged species such as **(S),** *either* the cation *or* the radical, can decompose by loss of hydrogen at C-3 to the 'normal' product **(2)** (path *a)* or by interception of the bridged species by a specific hydroxy group derived from the a-ketoglutarate-penicillin coupled reduction of dioxygen, which produced **(8)** (path *b).* Those processes emanating from the bridged cation would be heterolytic whereas the bridged radical would proceed through homolytic reactions.

 $\ddagger$  This preparation was shown to contain both DAOC synthetase and DAC synthetase activities by its ability, in separate experiments to convert both **(la)** into **(2)** and **(2)** into **(3);** see J. E. Baldwin, R. M. Adlington, J. R. Coates, M. J. C. Crabbe, N. P. Crouch, J. W. Keeping, G. C. Knight, C. J. Schofield, H.-H. Ting, C. A. Vallejo, M. Thorniley, and E. P. Abraham, *Biochem. J.,* 1987, **245,** 831.

<sup>§</sup> Partially purified DAOC/DAC synthetase (2 ml; *ca.* 0.5 International Units) in Tris-HC1 buffer (pH 7.4; 50 mM) was pre-incubated for *5* min at 27 "C and 250 rev. min-l with 200 **pl** of co-factor solution prepared from  $\alpha$ -ketoglutarate (14.6 mg), L-ascorbate (17.6 mg), dithiothreitol (30.8 mg), iron **(11)** sulphate (1.4 mg), and ammonium sulphate (1.32 g) in distilled water (10 ml). The substrate **(la)/(lb)** (1 mg) in Tris-HCl (pH 7.4; 1.8 ml; 50 mm) was added and the pH adjusted to 7.4 (NaOH). The resulting solution was incubated at  $27^{\circ}$ C, and  $250$  rev. min<sup>-1</sup> for 2 h, after which the protein was precipitated by the addition of acetone to 70% v/v. After centrifugation (10000 rev. min<sup>-1</sup>; 2 min;  $0^{\circ}$ C) the supernatant was evaporated to dryness and the residue dissolved in  $D<sub>2</sub>O$  (0.5 ml) and examined by n.m.r. (500 MHz;  $D<sub>2</sub>O$ ; HOD suppressed).

 $\int$  Prepared from  $[(R)-5-amino-5-carboxypentanov]$ -L-cysteinyl-p-[2-<sup>2</sup>H] valine by enzymic synthesis with isopenicillin N synthetase; see ref. 4. The level of deuteriation of C-3 was estimated to be  $> 98\%$  by 'H 500 MHz n.m.r. and mass spectral analysis.

<sup>\*\*</sup> Less than a quantitative incorporation of <sup>18</sup>O was expected as complete degassing of the enzyme solution (prepared in normal air) could not be achieved without extensive enzymic degradation.

tt Mixtures of **(la)** and **(lb)** are converted into **(2), (3),** and **(4)** with *no* isotopic enrichment in the pool of either **(la)** or **(lb).** Thus the isotope effect responsible for the changing ratio of  $(2) + (3)$  to  $(4)$ must occur subsequent to an irreversible step on a single enzyme, at a branching point in the reaction (see following paper).

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